

RAW SEQUENCE LISTING

DATE: 12/17/2001

PATENT APPLICATION: US/09/909,088

TIME: 15:26:15

Input Set : N:\Crf3\RULE60\09909088.txt

Output Set: N:\CRF3\12172001\I909088.raw

3 <110> APPLICANT: Genentech, Inc.
 4 Ashkenazi, Avi
 5 Botstein, David
 6 Desnoyers, Luc
 7 Eaton, Dan L.
 8 Ferrara, Napoleone
 9 Filvaroff, Ellen
 10 Fong, Sherman
 11 Gao, Wei-Qiang
 12 Gerber, Hanspeter
 13 Gerritsen, Mary E.
 14 Goddard, A.
 15 Godowski, Paul J.
 16 Grimaldi, Christopher J.
 17 Gurney, Austin L.
 18 Hillan, Kenneth, J.
 19 Kljavin, Ivar J.
 20 Mather, Jennie P.
 21 Pan, James
 22 Paoni, Nicholas F.
 23 Roy, Margaret Ann
 24 Stewart, Timothy A.
 25 Tumas, Daniel
 26 Williams, P. Mickey
 27 Wood, William, I.
 29 <120> TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
 30 Acids Encoding the Same
 32 <130> FILE REFERENCE: 10466-14
 34 <140> CURRENT APPLICATION NUMBER: 09/909,088
 35 <141> CURRENT FILING DATE: 2001-07-18
 37 <150> PRIOR APPLICATION NUMBER: 09/665,350
 38 <151> PRIOR FILING DATE: 2000-09-18
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 41 <151> PRIOR FILING DATE: 2000-02-22
 43 <150> PRIOR APPLICATION NUMBER: US 60/143,048
 44 <151> PRIOR FILING DATE: 1999-07-07
 46 <150> PRIOR APPLICATION NUMBER: US 60/145,698
 47 <151> PRIOR FILING DATE: 1999-07-26
 49 <150> PRIOR APPLICATION NUMBER: US 60/146,222
 50 <151> PRIOR FILING DATE: 1999-07-28
 52 <150> PRIOR APPLICATION NUMBER: PCT/US99/20594
 53 <151> PRIOR FILING DATE: 1999-09-08
 55 <150> PRIOR APPLICATION NUMBER: PCT/US99/20944
 56 <151> PRIOR FILING DATE: 1999-09-13
 58 <150> PRIOR APPLICATION NUMBER: PCT/US99/21090
 59 <151> PRIOR FILING DATE: 1999-09-15
 61 <150> PRIOR APPLICATION NUMBER: PCT/US99/21547

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104	tggagctccg	gctgctctt	cccgcagcgc	taccgcceat	gcgcctgccg	150
106	cgccgggccc	cgctggggct	cctgccgctt	ctgctgctgc	tgccgcccgc	200
108	gccggaggcc	gccaagaagc	cgacgccctg	ccaccgggtgc	cgggggctgg	250
110	tggacaagtt	taaccagggg	atggtggaca	ccgcaaagaa	gaactttggc	300
112	ggcgggaaca	cggtctggga	ggaaaagacg	ctgtccaagt	acgagtccag	350
114	cgagattcgc	ctgctggaga	tcctggaggg	gctgtgcgag	agcagcgact	400
116	tcgaatgcaa	tcagatgcta	gaggcgcaag	aggagcacct	ggaggcctgg	450
118	tggctgcagc	tgaagagcga	atatcctgac	ttattcgagt	ggttttgtgt	500
120	gaagacactg	aaagtgtgct	gctctccagg	aacctacggg	cccgaactgtc	550
122	tcgcattgcca	gggcgggatcc	cagaggccct	gcagcgggaa	tggccactgc	600
124	agcggagatg	ggagcagaca	gggcgacggg	tcctgccggg	gccacatggg	650
126	gtaccagggc	cgctgtgca	ctgactgcat	ggacggctac	ttcagctcgc	700
128	tccggaacga	gacccacagc	atctgcacag	cctgtgacga	gtcctgcaag	750
130	acgtgctcgg	gcctgaccaa	cagagactgc	ggcgagtgtg	aagtgggctg	800
132	ggtgctggac	gagggcgcct	gtgtggatgt	ggacgagtgt	gcggccgagc	850
134	cgctccctg	cagcgtgcg	cagttctgta	agaacgccaa	cggctcctac	900
136	acgtgcgaag	agtgtgactc	cagctgtgtg	ggctgcacag	gggaaggccc	950
138	aggaaactgt	aaagagtgtg	tctctggcta	cgcgaggagg	cacggacagt	1000
140	gtgcagatgt	ggacgagtgc	tcactagcag	aaaaaacctg	tgtgaggaaa	1050
142	aacgaaaact	gctacaatac	tccagggagc	tacgtctgtg	tgtgtcctga	1100
144	cggcttcgaa	gaaacggaag	atgcctgtgt	gccgccggca	gaggctgaag	1150
146	ccacagaag	agaaagcccc	acacagctgc	cctcccgcga	agacctgtaa	1200

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156  cttggttggt cttaaacaga cttgtatatt ttgatacagt tctttgtaat 1400
158  aaaattgacc attgtaggta atcaggagga aaaaaaaaaa aaaaaaaaaa 1450
160  aaagggcggc cgcgactcta gagtcgacct gcagaagctt ggccgccatg 1500
162  gcccaacttg tttattgcag cttataatgg ttacaaataa agcaatagca 1550
164  tcacaaatth cacaataaaa gcattttttt cactgcattc tagttgtggt 1600
166  ttgtccaaac tcatcaatgt atcttatcat gtctggatcg ggaattaatt 1650
168  cggcgcagca ccattggcctg aaataacctc tgaaagagga acttggttag 1700
170  gtaccttctg aggcggaaag aaccagctgt ggaatgtgtg tcagttaggg 1750
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174  ctcaattagt cagcaacca gtttt 1825
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186              20              25              30
188  Cys His Arg Cys Arg Gly Leu Val Asp Lys Phe Asn Gln Gly Met
189              35              40              45
191  Val Asp Thr Ala Lys Lys Asn Phe Gly Gly Gly Asn Thr Ala Trp
192              50              55              60
194  Glu Glu Lys Thr Leu Ser Lys Tyr Glu Ser Ser Glu Ile Arg Leu
195              65              70              75
197  Leu Glu Ile Leu Glu Gly Leu Cys Glu Ser Ser Asp Phe Glu Cys
198              80              85              90
200  Asn Gln Met Leu Glu Ala Gln Glu Glu His Leu Glu Ala Trp Trp
201              95              100             105
203  Leu Gln Leu Lys Ser Glu Tyr Pro Asp Leu Phe Glu Trp Phe Cys
204              110             115             120
206  Val Lys Thr Leu Lys Val Cys Cys Ser Pro Gly Thr Tyr Gly Pro
207              125             130             135
209  Asp Cys Leu Ala Cys Gln Gly Gly Ser Gln Arg Pro Cys Ser Gly
210              140             145             150
212  Asn Gly His Cys Ser Gly Asp Gly Ser Arg Gln Gly Asp Gly Ser
213              155             160             165
215  Cys Arg Cys His Met Gly Tyr Gln Gly Pro Leu Cys Thr Asp Cys
216              170             175             180
219  Met Asp Gly Tyr Phe Ser Ser Leu Arg Asn Glu Thr His Ser Ile
220              185             190             195
222  Cys Thr Ala Cys Asp Glu Ser Cys Lys Thr Cys Ser Gly Leu Thr
223              200             205             210
225  Asn Arg Asp Cys Gly Glu Cys Glu Val Gly Trp Val Leu Asp Glu
226              215             220             225
228  Gly Ala Cys Val Asp Val Asp Glu Cys Ala Ala Glu Pro Pro Pro

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Input Set : N:\Crf3\RULE60\09909088.txt

Output Set: N:\CRF3\12172001\I909088.raw

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234 Cys Glu Glu Cys Asp Ser Ser Cys Val Gly Cys Thr Gly Glu Gly
235          260          265          270
237 Pro Gly Asn Cys Lys Glu Cys Ile Ser Gly Tyr Ala Arg Glu His
238          275          280          285
240 Gly Gln Cys Ala Asp Val Asp Glu Cys Ser Leu Ala Glu Lys Thr
241          290          295          300
243 Cys Val Arg Lys Asn Glu Asn Cys Tyr Asn Thr Pro Gly Ser Tyr
244          305          310          315
246 Val Cys Val Cys Pro Asp Gly Phe Glu Glu Thr Glu Asp Ala Cys
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265 cgcccagccg tctaaacggg aacagccctg gctgagggag ctgcagcgca 150
267 gcagagtatc tgacggcgcc aggttgcgta ggtgcggcac gaggagtttt 200
269 cccggcagcg aggaggtcct gagcagcatg gcccgaggga gcgccttccc 250
271 tgccgcgcg ctctggctct ggagcatcct cctgtgcctg ctggcactgc 300
273 gggcgagggc cgggcgcgcg caggaggaga gcctgtacct atggatcgat 350
275 gctcaccagg caagagtaet cataggattt gaagaagata tcctgattgt 400
277 ttcagagggg aaaatggcac cttttacaca tgatttcaga aaagcgcaac 450
279 agagaatgcc agctattcct gtcaatatcc attccatgaa ttttacctgg 500
281 caagctgcag ggcaggcaga ataattctat gaattcctgt ccttgcgctc 550
284 cctggataaa ggcacatcag cagatccaac cgtcaatgtc cctctgctgg 600
286 gaacagtgcc tcacaaggca tcagttgttc aagttggttt cccatgtctt 650
288 ggaaaacagg atgggggtgc agcatttgaa gtggatgtga ttgttatgaa 700
290 ttctgaaggc aacaccattc tccaaacacc tcaaaatgct atcttcttta 750
292 aaacatgtca acaagctgag tgcccaggcg ggtgccgaaa tggaggcttt 800
294 tgtaatgaaa gacgcactcg cgagtgtcct gatgggttcc acggacctca 850
296 ctgtgagaaa gccctttgta ccccacgatg tatgaatggt ggactttgtg 900
298 tgactcctgg tttctgcata tgcccacctg gattctatgg agtgaactgt 950
300 gacaaagcaa actgtccaac cacctgcttt aatggaggga cctgtttcta 1000
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304 tcagcaaatg ccacacaacc tgcgaaatg gaggtaaatg cattggtaaa 1100
306 agcaaatgta agtgttccaa aggttaccag ggagacctct gttcaaaagc 1150
308 tgtctgcgag cctggctgtg gtgcacatgg aacctgccat gaacccaaca 1200
310 aatgccaatg tcaagaaggt tggcatggaa gacactgcaa taaaaggtag 1250
312 gaagccagcc tcatacatgc cctgaggcca gcaggcgccc agctcaggca 1300
314 gcacacgcct tacttataaa aggccgagga gcggcgggat ccacctgaat 1350

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Input Set : N:\Crif3\RULE60\09909088.txt

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316 ccaattacat ctggtgaact cgcacatctg aaacgtttta agttacacca 1400
318 agttcatagc ctttggttaac ctttcatgtg ttgaatgttc aaataatggt 1450
320 cattacactt aagaatactg gcctgaattt tattagcttc attataaate 1500
322 actgagctga tatctactct tccttttaag ttttctaagt acgtctgtag 1550
324 catgatggta tagattttct tgtttcagtg ctttgggaca gattttatat 1600
326 tatgtcaatt gatcagggtt aaattttcag tgtgtagtgt gcagatatatt 1650
328 tcaaaattac aatgcattta tgggtgtctg gggcagggga acatcagaaa 1700
330 gggttaaatt ggcaaaaaatg cgttaagtcac aagaatttgg atgggtgcagt 1750
332 taatgttgaa gttacagcat ttcagatttt attgtcagat atttagatgt 1800
334 ttgttacatt tttaaaaatt gctcttaatt tttaaactct caatacaata 1850
336 tattttgacc ttaccattat tccagagatt cagtattaaa aaaaaaaaaa 1900
338 ttacactgtg gtagtggcat ttaacaata taatatattc taaacacaat 1950
340 gaaataggga atataatgta tgaacttttt gcattggctt gaagcaatat 2000
342 aatataattg aaacaaaaca cagctcttac ctaataaaca ttttatactg 2050
344 tttgtatgta taaaataaag gtgctgcttt agttttttgg aaaaaaaaaa 2100
346 aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa gggcgccgcg gactctagag 2150
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365 Pro Gln Glu Glu Ser Leu Tyr Leu Trp Ile Asp Ala His Gln Ala
366 35 40 45
368 Arg Val Leu Ile Gly Phe Glu Glu Asp Ile Leu Ile Val Ser Glu
369 50 55 60
371 Gly Lys Met Ala Pro Phe Thr His Asp Phe Arg Lys Ala Gln Gln
372 65 70 75
374 Arg Met Pro Ala Ile Pro Val Asn Ile His Ser Met Asn Phe Thr
375 80 85 90
377 Trp Gln Ala Ala Gly Gln Ala Glu Tyr Phe Tyr Glu Phe Leu Ser
378 95 100 105
380 Leu Arg Ser Leu Asp Lys Gly Ile Met Ala Asp Pro Thr Val Asn
381 110 115 120
383 Val Pro Leu Leu Gly Thr Val Pro His Lys Ala Ser Val Val Gln
384 125 130 135
386 Val Gly Phe Pro Cys Leu Gly Lys Gln Asp Gly Val Ala Ala Phe
387 140 145 150
389 Glu Val Asp Val Ile Val Met Asn Ser Glu Gly Asn Thr Ile Leu
390 155 160 165
392 Gln Thr Pro Gln Asn Ala Ile Phe Phe Lys Thr Cys Gln Gln Ala
393 170 175 180
395 Glu Cys Pro Gly Gly Cys Arg Asn Gly Gly Phe Cys Asn Glu Arg
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VERIFICATION SUMMARY

PATENT APPLICATION: US/09/909,088

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